Access DB# 99583

# SEARCH REQUEST FORM

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Please provide a detailed statement of the search topic. Include the elected species or structures, keywords, synutility of the invention. Define any terms that may have known. Please attach a copy of the cover sheet, pertiner	onyms, acronyms, and regis e a special meaning. Give e	try numbers, and combine with the concept or	ı
Title of Invention: Tissue Remode Inventors (please provide full names): S. Be			
Earliest Priority Filing Date: [2 -31 - 200]		į.	
*For Sequence Searches Only* Please include all pertinen appropriate serial number.		ivisional, or issued patent numbers) along with the	
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FILE COVERS 1907 - 28 Jul 2003 VOL 139 ISS 5 FILE LAST UPDATED: 27 Jul 2003 (20030727/ED)

This file contains CAS Registry Numbers for easy and accurate substance identification.

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          42629 SEA FILE=REGISTRY ABB=ON PLU=ON GGIV|IVEE|VEEY|EEYQ|EYQL|YQLP
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            339 SEA FILE=REGISTRY ABB=ON
                                          PLU=ON L1 AND SQL<=25
L4
          50431 SEA FILE=REGISTRY ABB=ON PLU=ON
                                                 BONE OR FATTY(W) ACID? OR
                ACYLAT?
         686265 SEA FILE=HCAPLUS ABB=ON PLU=ON L4 OR BONE OR FATTY(W)ACID?
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                OR ACYLAT? OR OSTEOPOROS?
L10
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L12
             20 SEA FILE=HCAPLUS ABB=ON
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L12 ANSWER 1 OF 20 HCAPLUS COPYRIGHT 2003 ACS on STN

ACCESSION NUMBER:

2003:306849 HCAPLUS

DOCUMENT NUMBER:

138:332694

TITLE:

Polymorphism identification within 50 equine gene-specific sequence tagged sites. [Erratum to

document cited in CA136:211522]

AUTHOR(S):

Shubitowski, D. M.; Venta, P. J.; Douglass, C. L.;

Zhou, R.-X.; Weart, S. L.

CORPORATE SOURCE:

Department of Large Animal Clinical Sciences, Michigan

State University, East Lansing, MI, 48824, USA

SOURCE: Animal Genetics (2001), 32(5), 332

CODEN: ANGEE3; ISSN: 0268-9146

PUBLISHER:

Blackwell Science Ltd.

DOCUMENT TYPE:

Journal

LANGUAGE:

English

One sequence (GenBank AY008814) was misidentified as being part of the equine sex-detg. region gene (SRY). The correct identity of the sequence is as part of the sex-detg. region Y box 30 gene (SOX30).

ΙT 402741-21-3 402809-83-0

#### Russel 10 032330

RL: BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study) (amino acid sequence; polymorphism identification within 50 equine gene-specific sequence tagged sites (Erratum)) L12 ANSWER 2 OF 20 HCAPLUS COPYRIGHT 2003 ACS on STN ACCESSION NUMBER: 2002:850307 HCAPLUS DOCUMENT NUMBER: 137:346244 TITLE: Tissue remodeling with compds. comprising a sequence from TGF-.beta. super family Ser/Thr/kinase receptors INVENTOR(S): Ben-Sasson, Shmuel PATENT ASSIGNEE(S): Children's Medical Center Corporation, USA U.S. Pat. Appl. Publ., 51 pp., Cont.-in-part of Appl. SOURCE: No. PCT/US00/32852. CODEN: USXXCO DOCUMENT TYPE: Patent LANGUAGE: English FAMILY ACC. NUM. COUNT: PATENT INFORMATION: PATENT NO. KIND DATE APPLICATION NO. US 2002165150 A120021107 US 2001-32330 20011231 WO 2001042280 Α2 20010614 WO 2000-US32852 20001204 WO 2001042280 А3 20020307 W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG PRIORITY APPLN. INFO.: US 1998-161094 B2 19980925 WO 2000-US32851 W 20001204 WO 2000-US32852 A2 20001204 US 1999-458491 A1 19991209 The invention concerns a method for the modulation of tissue-remodeling processes, by contacting the tissue to be remodeled with a compd. comprising a sequence derived from certain regions of TGF-.beta. super family Ser/Thr/kinase receptors. **332350-87-5,** BMP receptor kinase-2 RL: BSU (Biological study, unclassified); BIOL (Biological study) (tissue remodeling with compds. comprising a sequence from TGF-.beta. super family Ser/Thr/kinase receptors) 474526-71-1 474526-72-2 474526-73-3 474526-78-8 474526-96-0 474526-98-2 474527-06-5 RL: PAC (Pharmacological activity); THU (Therapeutic use); BIOL (Biological study); USES (Uses) (tissue remodeling with compds. comprising a sequence from TGF-.beta. super family Ser/Thr/kinase receptors) L12 ANSWER 3 OF 20 HCAPLUS COPYRIGHT 2003 ACS on STN 2002:691570 HCAPLUS ACCESSION NUMBER: DOCUMENT NUMBER: 137:196722 TITLE: cDNA and protein sequence of a novel human actin related-protein ARP sequence homolog protein 32 and their uses in drug screening, diagnosis and therapeutics

AB

IT

IT

INVENTOR(S):

PATENT ASSIGNEE(S):

Bode Gene Development Co., Ltd., Shanghai, Peop. Rep.

Mao, Yumin; Xie, Yi

## Russel 10 032330

China

SOURCE:

Faming Zhuanli Shenqing Gongkai Shuomingshu, 36 pp.

CODEN: CNXXEV

DOCUMENT TYPE: LANGUAGE:

Patent Chinese

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO. KIND DATE APPLICATION NO. DATE

CN 1328038 A 20011226 CN 2000-116455 20000612

PRIORITY APPLN. INFO.: CN 2000-116455 20000612

This invention provides the cDNA and protein sequence of a novel human actin related-protein ARP sequence homolog protein 32 cloned from fetal brain. The mol. wt. of protein 32 is 32 kDa detd. on SDS PAGE and the sequence of protein 32 has homol. with that of actin related-protein ARP. The invention discloses the process of screening the agonist and antagonist against the polypeptide. The protein 32 can be used to diagnosis and treatment for many actin related-protein ARP assocd. diseases such as blood diseases, bone and vessel development disorders, kidney diseases, inflammation and immune diseases.

IT 452093-05-9

RL: PRP (Properties)

(unclaimed sequence; cDNA and protein sequence of a novel human actin related-protein ARP sequence homolog protein 32 and their uses in drug screening, diagnosis and therapeutics)

L12 ANSWER 4 OF 20 HCAPLUS COPYRIGHT 2003 ACS on STN

ACCESSION NUMBER:

2002:575193 HCAPLUS

DOCUMENT NUMBER:

137:139362

TITLE:

Human antibodies and fragments derived from phage

display library for selective cancer therapy and

diagnosis

INVENTOR(S):

Hagai, Yocheved; Lazarovits, Janette; Guy, Rachel; Lipschitz, Orly; Szanton, Esther; Levanon, Avigdor;

Plaksin, Daniel; Peretz, Tuvia Bio-Technology General Corp., USA

PATENT ASSIGNEE(S):

PCT Int. Appl., 232 pp.

SOURCE:

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT: 2

PATENT INFORMATION:

	PATENT NO.			KI	ND	DATE			A	PPLI	CATI	ON N	ο.	DATE					
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and/or specifically to a target cell in favor of other cells, wherein the

#### Russel 10 032330

binding selectivity or specificity is primarily detd. by a first hypervariable region, and wherein the Fv is a scFv or a dsFv, and optionally having one or more tags. The enhanced binding is directed to a substantially exposed and/or over-expressed binding site on or in a target comprising a cell in favor of other cells on or in which the binding site is not substantially available and/or expressed. The invention is further directed to a method for isolating such peptides and polypeptides from a phage display library and to the nucleic acid mols. encoding them. The invention provides for a pharmaceutical compn. comprising the peptide or polypeptide and kits for diagnosis and treatment of disease, specifically cancer, most specifically acute myeloid leukemia.

#### IT 442527-58-4P

RL: BPN (Biosynthetic preparation); BSU (Biological study, unclassified); DGN (Diagnostic use); PRP (Properties); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); USES (Uses)

(human antibodies and fragments derived from phage display library for selective cancer therapy and diagnosis)

L12 ANSWER 5 OF 20 HCAPLUS COPYRIGHT 2003 ACS on STN

ACCESSION NUMBER:

2002:382316 HCAPLUS

DOCUMENT NUMBER:

137:16533

TITLE:

Human genome-derived single exon nucleic acid probes useful for analysis of gene expression in human lung Penn, Sharron G.; Hanzel, David K.; Chen, Wensheng;

INVENTOR(S):

Rank, David R.

Kalik

PATENT ASSIGNEE(S):

Molecular Dynamics, Inc., USA

SOURCE:

PCT Int. Appl., 634 pp.

CODEN: PIXXD2

DOCUMENT TYPE: LANGUAGE:

Patent English

FAMILY ACC. NUM. COUNT:

89

PATENT INFORMATION:

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GB 2000-24263
                 Α
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WO 2001-US665
                 W
                     20010130
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AΒ A single exon nucleic acid microarray comprising 12,614 single exon nucleic acid probes for measuring gene expression in a sample derived from human lung cells is described. These unique exons are within longer probe sequences; sequencing confirms the exact chem. structure of each probe. Some amplicons have more than one exon, and some exons are contained in more than one amplicon. Expression, homol., and functional information are provided for the genome-derived single exon probes that are expressed significantly in human lung. Also described are 12,386 single exon nucleic acid probes and 12,011 proteins expressed in the lung and their use in methods for detecting gene expression. The genome-derived single exon nucleic acids comprise a novel type of nucleic acid microarray for verifying gene expression. In addn., methods are provided for identifying exons in a eukaryotic genome, and for assigning exons to a single gene. [This abstr. record is one of nine records for this document necessitated by the large no. of index entries required to fully index the document and

by the large no. of index entries r publication system constraints.].

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400618-81-7
RL: BSU (Biological study, unclassified); PRP (Properties); BIOL
(Biological study)
   (amino acid sequence; human genome-derived single exon nucleic acid
  probes useful for anal. of gene expression in human lung)
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PATENT ASSIGNEE(S):
                         USA
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AB Methods and app. for predicting, confirming and displaying functional regions from genomic sequence data are used to identify 16,834 unique human genome-derived single exon probes useful for gene expression anal., particularly gene expression anal. by microarray. Also presented are genome-derived single exon microarrays that include such probes, peptides encoded by the exons, and antibodies thereto. The human genome-derived single-exon probes are known to be expressed in one or more human tissues or cell types, particularly human brain, heart, liver, fetal liver, placenta, lung, bone marrow, BT474 and other human mammary epithelial cells, HeLa and other human cervical epithelial cells, and HBL 100 and other human mammary epithelial cells. The invention provides a method of financing, selling and/or licensing genome-derived single-exon microarrays to customer desiring to measure gene expression, comprising: making available for computerized query or subscription service a database having a record corresponding to each genome-derived single exon microarray available for sale and/or license. [This abstr. record is one of ten records for this document necessitated by the large no. of index entries required to fully index the document and publication system constraints.].

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INVENTOR(S):
                         Penn, Sharron G.; Hanzel, David K.; Chen, Wensheng;
                         Rank, David R.
PATENT ASSIGNEE(S):
                         Molecular Dynamics, Inc., USA
SOURCE:
                         PCT Int. Appl., 658 pp.
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DOCUMENT TYPE:
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AB
     A single exon nucleic acid microarray comprising 13,109 single exon
     nucleic acid probes for measuring gene expression in a sample derived from
     human adult liver is described. These unique exons are within longer
     probe sequences; sequencing confirms the exact chem. structure of each
     probe. Some amplicons have more than one exon, and some exons are
     contained in more than one amplicon. Expression, homol., and functional
     information are provided for the genome-derived single exon probes that
    are expressed significantly in human adult liver cells. Also described
    are 12,886 single exon nucleic acid probes and 12,583 proteins expressed
    in the adult liver and their use in methods for detecting gene expression.
    The genome-derived single exon nucleic acids comprise a novel type of
    nucleic acid microarray for verifying gene expression. In addn., methods
    are provided for identifying exons in a eukaryotic genome, and for
    assigning exons to a single gene.
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ACCESSION NUMBER:
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DOCUMENT NUMBER:
                          136:215388
TITLE:
                          Immunogenic hepatitis B nucleocapsid protein (HBc)
                          chimeric particles having enhanced stability
INVENTOR(S):
                          Birkett, Ashley J.
PATENT ASSIGNEE(S):
                          Apovia, Inc., USA
SOURCE:
                          PCT Int. Appl., 290 pp.
                          CODEN: PIXXD2
DOCUMENT TYPE:
                          Patent
LANGUAGE:
                          English
FAMILY ACC. NUM. COUNT:
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                                         WO 2001-US41759 W 20010816
    A chimeric, carboxy-terminal truncated hepatitis B virus nucleocapsid
    protein (core protein or HBc) is disclosed that is engineered for both
     enhanced stability of self-assembled particles and the display of an
     immunogenic epitope. The immunogenic epitope is a B cell epitope or T
     cell epitope derived from pathogen such as Streptococcus pneumonia,
    Cryptosporidium parvum, HIV, foot and mouth disease virus, influenza
    virus, Yersinia pestia, etc. The display of the immunogenic epitope is
    displayed in the immunogenic loop of HBc, whereas the enhanced stability
     of self-assembled particles is obtained by the presence of at least one
    heterologous cysteine residue near the carboxy-terminus of the chimer mol.
    Methods of making and using the chimers are also disclosed.
    112-80-1D, Oleic acid, sorbitan or mannitol esters
    RL: BSU (Biological study, unclassified); THU (Therapeutic use); BIOL
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        (chimeric proteins comprising HBcAg and T and/or B cell epitope for use
        as vaccines)
    401460-88-6
    RL: PRP (Properties)
        (unclaimed sequence; immunogenic hepatitis B nucleocapsid protein (HBc)
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AB

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chimeric particles having enhanced stability)

## Russel 10 032330

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L12 ANSWER 9 OF 20
                         HCAPLUS COPYRIGHT 2003 ACS on STN
ACCESSION NUMBER:
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DOCUMENT NUMBER:
                              136:195269
TITLE:
                             Human genome-derived single exon nucleic acid probes
                              useful for analysis of gene expression in human
                             placenta
INVENTOR(S):
                             Penn, Sharron G.; Hanzel, David K.; Chen, Wensheng;
                             Rank, David R.
PATENT ASSIGNEE(S):
                             Molecular Dynamics, Inc., USA
SOURCE:
                             PCT Int. Appl., 654 pp.
                             CODEN: PIXXD2
DOCUMENT TYPE:
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     A single exon nucleic acid microarray comprising 13,232 single exon
AB
     nucleic acid probes for measuring gene expression in a sample derived from
     human placenta cells is described. These unique exons are within longer
     probe sequences; sequencing confirms the exact chem. structure of each
     probe. Some amplicons have more than one exon, and some exons are
     contained in more than one amplicon. Expression, homol., and functional
     information are provided for the genome-derived single exon probes that
     are expressed significantly in human placenta. Also described are 13,000
     single exon nucleic acid probes and 12,605 proteins expressed in the
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placenta cells and their use in methods for detecting gene expression. The genome-derived single exon nucleic acids comprise a novel type of

nucleic acid microarray for verifying gene expression. In addn., methods

are provided for identifying exons in a eukaryotic genome, and for assigning exons to a single gene. [This abstr. record is one of nine records for this document necessitated by the large no. of index entries required to fully index the document and publication system constraints.]. 388091-67-6 388091-68-7 388091-69-8 388091-71-2:388091-72-3 388091-73-4 388091-74-5 388091-76-7 388091-77-8 388091-78-9 388091-79-0 388091-80-3 388091-82-5 388091-83-6 388091-84-7 388091-85-8 388091-86-9 388091-87-0 388091-89-2 388091-90-5 388091-91-6 388091-92-7 388091-93-8 388091-94-9 388091-95-0 388091-96-1 388091-97-2 388091-98-3 388091-99-4 388092-00-0 388092-01-1 388092-02-2 388092-03-3 388092-04-4 388092-05-5 388092-06-6 388092-07-7 388092-08-8 388092-09-9 388092-10-2 388092-11-3 388092-12-4 388092-13-5 388092-14-6 388092-15-7 400616-76-4 400616-77-5 400616-78-6 400616-79-7 400616-80-0 400616-81-1 400616-82-2 400616-85-5 400616-86-6 400616-87-7 400616-88-8 400616-89-9 400616-90-2 400616-91-3 400616-92-4 400616-93-5 400616-94-6 400616-95-7 400616-96-8 400616-97-9 400616-98-0 400616-99-1 400617-00-7 400617-01-8 400617-02-9 400617-03-0 400617-04-1 400617-05-2 400617-06-3 400617-07-4 400617-08-5 400617-09-6 400617-10-9 400617-11-0 400617-12-1 400617-13-2 400617-14-3 400617-15-4 400617-16-5 400617-17-6 400617-18-7 400617-19-8 400617-20-1 400617-21-2 400617-22-3 400617-23-4 400617-24-5 400617-25-6 400617-26-7 400617-27-8 400617-28-9 400617-29-0 400617-30-3 400617-31-4 400617-32-5 400617-33-6 400617-34-7 400617-35-8 400617-36-9 400617-37-0 400617-38-1 400617-39-2 400617-40-5 400617-41-6 400617-43-8 400617-44-9 400617-45-0 400617-46-1 400617-47-2 400617-48-3 400617-49-4 400617-50-7 400617-51-8 400617-52-9 400617-53-0 400617-54-1 400617-55-2 400617-56-3 400617-57-4 400617-58-5 400617-59-6 400617-60-9 400617-61-0 400617-62-1 400617-63-2 400617-64-3 400617-65-4 400617-66-5 400617-67-6 400617-68-7 400617-69-8 400617-70-1 400617-71-2 400617-72-3 400617-73-4 400617-74-5 400617-75-6 400617-76-7 400617-77-8 400617-78-9 400617-79-0 400617-80-3 400617-81-4 400617-82-5 400617-83-6 400617-84-7 400617-86-9 400617-87-0 400617-88-1 400617-89-2 400617-90-5 400617-91-6 400617-92-7 400617-93-8 400617-94-9 400617-95-0 400617-96-1 400617-97-2 400617-98-3 400618-00-0 400618-01-1 400618-02-2 400618-03-3 400618-04-4 400618-05-5 400618-06-6

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RL: ANT (Analyte); BSU (Biological study, unclassified); PRP (Properties);
ANST (Analytical study); BIOL (Biological study)
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## Russel 10\_032330

(amino acid sequence; human genome-derived single exon nucleic acid probes useful for anal. of gene expression in human placenta) ΙT 400624-90-0 400624-91-1 400624-92-2 400624-94-4 400624-95-5 400624-96-6 400624-97-7 400624-98-8 400624-99-9 400625-01-6 400625-02-7 400631-91-6 400632-33-9 400632-35-1 400632-37-3 400632-38-4 400632-40-8 400632-43-1 400632-44-2 400632-46-4 400632-47-5 400632-49-7 400632-50-0 400632-52-2 400632-54-4 400632-55-5 400632-58-8 400632-59-9 400632-62-4 400632-63-5 400632-66-8 400632-67-9 400632-69-1 400632-70-4 400632-72-6 400632-73-7 400632-74-8 400632-77-1 400632-78-2 RL: ANT (Analyte); BSU (Biological study, unclassified); PRP (Properties); ANST (Analytical study); BIOL (Biological study) (amino acid sequence; human genome-derived single exon nucleic acid probes useful for anal. of gene expression in human placenta)

L12 ANSWER 10 OF 20 HCAPLUS COPYRIGHT 2003 ACS on STN

ACCESSION NUMBER: DOCUMENT NUMBER:

2002:110610 HCAPLUS

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136:351347

TITLE:

Human genome-derived single exon nucleic acid probes useful for analysis of gene expression in human HeLa

cells or other human cervical epithelial cells

Penn, Sharron G.; Hanzel, David K.; Chen, Wensheng;

Rank, David R.

PATENT ASSIGNEE(S):

Molecular Dynamics, Inc., USA

SOURCE:

PCT Int. Appl., 487 pp. CODEN: PIXXD2

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DOCUMENT TYPE:

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LANGUAGE:

INVENTOR(S):

English

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PATENT INFORMATION:

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AΒ
     A single exon nucleic acid microarray comprising 9290 single exon nucleic
     acid probes for measuring gene expression in a sample derived from human
     HeLa cells or other human cervical epithelial cells is described. These
     unique exons are within longer probe sequences; sequencing confirms the
     exact chem. structure of each probe. Some amplicons have more than one
     exon, and some exons are contained in more than one amplicon. Expression,
     homol., and functional information are provided for the genome-derived
     single exon probes that are expressed significantly in human HeLa cells or
     other human cervical epithelial cell lines. Also described are 9102
     single exon nucleic acid probes and 8549 proteins expressed in the
     cervical epithelial cells and their use in methods for detecting gene
     expression. The genome-derived single exon nucleic acids comprise a novel
     type of nucleic acid microarray for verifying gene expression. In addn.,
    methods are provided for identifying exons in a eukaryotic genome, and for
     assigning exons to a single gene. [This abstr. record is one of six
     records for this document necessitated by the large no. of index entries
     required to fully index the document and publication system constraints.].
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ACCESSION NUMBER:
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TITLE: ·
                         Human genome-derived single exon nucleic acid probes
                         useful for analysis of gene expression in human fetal
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                         Penn, Sharron G.; Hanzel, David K.; Chen, Wensheng;
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INVENTOR(S):

Rank, David R.

PATENT ASSIGNEE(S):

Molecular Dynamics, Inc., USA

PCT Int. Appl., 639 pp.

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89

PATENT INFORMATION:

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     A single exon nucleic acid microarray comprising 12,673 single exon
AB
     nucleic acid probes for measuring gene expression in a sample derived from
     human fetal liver cells is described. These unique exons are within
     longer probe sequences; sequencing confirms the exact chem. structure of
     each probe. Some amplicons have more than one exon, and some exons are
     contained in more than one amplicon. Expression, homol., and functional
     information are provided for the genome-derived single exon probes that
     are expressed significantly in human fetal liver cells. Also described
     are 12,456 single exon nucleic acid probes and 12,027 proteins expressed
     in the fetal liver and their use in methods for detecting gene expression.
     The genome-derived single exon nucleic acids comprise a novel type of
     nucleic acid microarray for verifying gene expression. In addn., methods
     are provided for identifying exons in a eukaryotic genome, and for
     assigning exons to a single gene. [This abstr. record is one of nine
     records for this document necessitated by the large no. of index entries
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                         Penn, Sharron G.; Hanzel, David K.; Chen, Wensheng;
                         Rank, David R.
PATENT ASSIGNEE(S):
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A single exon nucleic acid microarray comprising 12,821 single exon nucleic acid probes for measuring gene expression in a sample derived from human brain cells is described. These unique exons are within longer probe sequences; sequencing confirms the exact chem. structure of each probe. Some amplicons have more than one exon, and some exons are contained in more than one amplicon. Expression, homol., and functional information are provided for the genome-derived single exon probes that are expressed significantly in human brain. Also described are 12,613 single exon nucleic acid probes and 12,377 proteins expressed in the brain and their use in methods for detecting gene expression. The genome-derived single exon nucleic acids comprise a novel type of nucleic acid microarray for verifying gene expression. In addn., methods are provided for identifying exons in a eukaryotic genome, and for assigning exons to a single gene. [This abstr. record is one of nine records for this document necessitated by the large no. of index entries required to fully index the document and publication system constraints.].

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        probes useful for anal. of gene expression in human brain)
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DOCUMENT NUMBER:
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TITLE:
                          Human genome-derived single exon nucleic acid probes
                          useful for analysis of gene expression in human bone
                          marrow
INVENTOR(S):
                          Penn, Sharron G.; Hanzel, David K.; Chen, Wensheng;
                          Rank, David R.
PATENT ASSIGNEE(S):
                          Molecular Dynamics, Inc., USA
SOURCE:
                          PCT Int. Appl., 657 pp.
                          CODEN: PIXXD2
DOCUMENT TYPE:
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LANGUAGE:
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FAMILY ACC. NUM. COUNT:
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AB
     A single exon nucleic acid microarray comprising 13,114 single exon
     nucleic acid probes for measuring gene expression in a sample derived from
     human bone marrow is described. These unique exons are within longer
     probe sequences; sequencing confirms the exact chem. structure of each
     probe. Some amplicons have more than one exon, and some exons are
     contained in more than one amplicon. Expression, homol., and functional
     information are provided for the genome-derived single exon probes that
     are expressed significantly in human bone marrow. Also described are
     12,898 single exon nucleic acid probes and 12,616 proteins expressed in
     the bone marrow and their use in methods for detecting gene expression.
     The genome-derived single exon nucleic acids comprise a novel type of
     nucleic acid microarray for verifying gene expression. In addn., methods
     are provided for identifying exons in a eukaryotic genome, and for
     assigning exons to a single gene. [This abstr. record is one of nine
     records for this document necessitated by the large no. of index entries
     required to fully index the document and publication system constraints.].
L12 ANSWER 14 OF 20 HCAPLUS COPYRIGHT 2003 ACS on STN
ACCESSION NUMBER:
                          2002:64569 HCAPLUS
DOCUMENT NUMBER:
                          136:195264
TITLE:
                         Human genome-derived single exon nucleic acid probes
                         useful for analysis of gene expression in human heart
INVENTOR(S):
                         Penn, Sharron G.; Hanzel, David K.; Chen, Wensheng;
                         Rank, David R.
PATENT ASSIGNEE(S):
                         Molecular Dynamics, Inc., USA
                         PCT Int. Appl., 529 pp.
SOURCE:
                         CODEN: PIXXD2
DOCUMENT TYPE:
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LANGUAGE:
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FAMILY ACC. NUM. COUNT:
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     A single exon nucleic acid microarray comprising 9980 single exon nucleic
AB
     acid probes for measuring gene expression in a sample derived from human
     heart is described. These unique exons are within longer probe sequences;
     sequencing confirms the exact chem. structure of each probe. Some
     amplicons have more than one exon, and some exons are contained in more
     than one amplicon. Expression, homol., and functional information are
     provided for the genome-derived single exon probes that are expressed
     significantly in human heart cells. Also described are 9791 single exon
     nucleic acid probes and 9347 proteins expressed in the heart and their use
     in methods for detecting gene expression. The genome-derived single exon
     nucleic acids comprise a novel type of nucleic acid microarray for
     verifying gene expression. In addn., methods are provided for identifying
     exons in a eukaryotic genome, and for assigning exons to a single gene.
     [This abstr. record is one of six records for this document necessitated
     by the large no. of index entries required to fully index the document and
     publication system constraints.].
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TITLE:

Human genome-derived single exon nucleic acid probes useful for analysis of gene expression in human breast

and BT 474 cells

INVENTOR(S):

Penn, Sharron G.; Hanzel, David K.; Chen, Wensheng;

Rank, David R.

PATENT ASSIGNEE(S):

Molecular Dynamics, Inc., USA

SOURCE:

PCT Int. Appl., 327 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent English

LANGUAGE: FAMILY ACC. NUM. COUNT:

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PATENT INFORMATION:

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AB
     A single exon nucleic acid microarray comprising 5205 single exon nucleic
     BT 474 cells is described. These unique exons are within longer probe
     sequences; sequencing confirms the exact chem. structure of each probe.
     Some amplicons have more than one exon, and some exons are contained in
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acid probes for measuring gene expression in a sample derived from human BT 474 cells is described. These unique exons are within longer probe sequences; sequencing confirms the exact chem. structure of each probe. Some amplicons have more than one exon, and some exons are contained in more than one amplicon. Expression, homol., and functional information are provided for the genome-derived single exon probes that are expressed significantly in human BT 474 cells. Also described are 5112 single exon nucleic acid probes and 5121 proteins expressed in the BT 474 cells and their use in methods for detecting gene expression. The genome-derived single exon nucleic acids comprise a novel type of nucleic acid microarray for verifying gene expression. In addn., methods are provided for identifying exons in a eukaryotic genome, and for assigning exons to a single gene. [This abstr. record is one of three records for this document necessitated by the large no. of index entries required to fully index the document and publication system constraints.].

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400625-15-2
RL: BSU (Biological study, unclassified); PRP (Properties); BIOL
(Biological study)
   (amino acid sequence; human genome-derived single exon nucleic acid
  probes useful for anal. of gene expression in human breast and BT 474
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     RL: BSU (Biological study, unclassified); PRP (Properties); BIOL
     (Biological study)
        (amino acid sequence; human genome-derived single exon nucleic acid
        probes useful for anal. of gene expression in human breast and BT 474
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     RL: BSU (Biological study, unclassified); PRP (Properties); BIOL
     (Biological study)
        (amino acid sequence; human genome-derived single exon nucleic acid
        probes useful for anal. of gene expression in human breast and BT 474
        cells)
L12 ANSWER 16 OF 20
                      HCAPLUS COPYRIGHT 2003 ACS on STN
ACCESSION NUMBER:
                         2001:512017
                                      HCAPLUS
DOCUMENT NUMBER:
                         136:211522
                         Polymorphism identification within 50 equine
TITLE:
                         gene-specific sequence tagged sites
AUTHOR(S):
                         Shubitowski, D. M.; Venta, P. J.; Douglass, C. L.;
                         Zhou, R.-X.; Ewart, S. L.
CORPORATE SOURCE:
                         Department of Large Animal Clinical Sciences, Michigan
                         State University, East Lansing, MI, 48824, USA Animal Genetics (2001), 32(2), 78-88
SOURCE:
                         CODEN: ANGEE3; ISSN: 0268-9146
PUBLISHER:
                         Blackwell Science Ltd.
DOCUMENT TYPE:
                         Journal
                         English
LANGUAGE:
     The continued discovery of polymorphisms in the equine genome will be
     important for future studies using genomic screens and fine mapping for
     the identification of disease genes. Segments of 50 equine genes were
     examd. for variability in 10 different horse breeds using a
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pool-and-sequence method. We identified 11 single nucleotide

polymorphisms (SNPs) in 9380 bp of sequenced exon, and 25 SNPs, six microsatellites, and one insertion/deletion in 16961 bp of sequenced intron. Of all genes studied 52% contained at least one polymorphism, and polymorphisms were found at an overall rate of 1/613 bp. Several of the putative SNPs were tested and verified by restriction enzyme anal. using natural restriction sites or ones created by primer mutagenesis. The lowest allele frequency for a SNP detected in pooled samples was 10%. Three of the SNPs verified in the diverse horse pool were further tested in six breed-specific horse pools and were found to be reasonably variable within breeds. The pool-and-sequence method allows identification of polymorphisms in horse populations and will be a valuable tool for future disease gene and comparative mapping in horses.

# IT 402741-21-3 402809-83-0

RL: BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study)

(amino acid sequence; polymorphism identification within 50 equine gene-specific sequence tagged sites)

REFERENCE COUNT:

THERE ARE 26 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L12 ANSWER 17 OF 20 HCAPLUS COPYRIGHT 2003 ACS on STN

ACCESSION NUMBER:

2000:34995 HCAPLUS

DOCUMENT NUMBER:

132:102856

TITLE:

Hyaluronic acid mimics for treatment of inflammation

and other hyaluronate-associated diseases

INVENTOR(S):

Prestwich, Glenn D.; Ziebell, Michael; Luo, Bai; Zhao,

Zhan-Gong

PATENT ASSIGNEE(S):

USA

SOURCE:

PCT Int. Appl., 53 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

P	PATENT NO.			KI	ND	DATE			A	PPLI	CATI	ON N	ο.	DATE				
	WO 2000001841 WO 2000001841								WO 1999-US15263					19990706				
"									BB,	BG,	BR,	BY,	CA,	CH,	CN,	CU,	CZ,	DE,
			DK,	EE,	ES,	FI,	GB,	GE,	GH,	GM,	HR,	HU,	ID,	IL,	IS,	JP,	KE,	KG,
															MK,			
			NO,	NZ,	PL,	PT,	RO,	RU,	SD,	SE,	SG,	SI,	SK,	SL,	ТJ,	TM,	TR,	TT,
			UA,	UG,	US,	UZ,	VN,	YU,	ZW,	AM,	ΑZ,	BY,	KG,	ΚZ,	MD,	RU,	ТJ,	TM
		RW:	GH,	GM,	ΚE,	LS,	MW,	SD,	SL,	SZ,	UG,	ZW,	AT,	·BE,	CH,	CY,	DE,	DK,
			ES,	FΙ,	FR,	GB,	GR,	IE,	IT,	LU,	MC,	NL,	PT,	SE,	BF,	ВJ,	CF,	CG,
			CI,	CM,	GA,	GN,	GW,	ML,	MR,	NE,	SN,	TD,	TG					
С	Ά	2346	742		A	A	2000	0113		C.	A 19	99-2	3467	42	1999	0706		
A	ΔU	9949	716		Α	1 .	2000	0124		Α	U 19:	99-4	9716		1999	0706		
. E	P.	11690	048		A.	2	2002	0109		E	P 19	99-9	337:1	8 '	1999	0706		
		R:	AT,	BE,	CH,	DE,	DK,	ES,	FR,	GB,	GR,	IT,	LI,	LU,	NL,	SE,	MC,	PT,
			ΙE,	SI,	LT,	LV,	FI,	RO										
PRIORI	PRIORITY APPLN. INFO.:						1	US 1	998-	9175	8P	P	1998	0706				
									1	US 1	999-	3477	07	Α	1999	0703		
				•					1	WO 1	999-1	US15:	263	W	1999	0706		
AB H	AB HA mimics and methods related thereto are disclosed. In particular,																	

AB HA mimics and methods related thereto are disclosed. In particular, mimics with structures detd. by virtue of novel methods, and the novel methods are disclosed. The HA mimics are useful for a variety of HA-related uses, including treatment of inflammatory diseases, tumor angiogenesis, skin disease, bone disease, and cardiovascular diseases.

### IT 254965-50-9P

RL: BAC (Biological activity or effector, except adverse); BPR (Biological

process); BSU (Biological study, unclassified); PNU (Preparation,
unclassified); PRP (Properties); THU (Therapeutic use); BIOL (Biological
study); PREP (Preparation); PROC (Process); USES (Uses)
 (hyaluronic acid mimics for treatment of inflammation and other
 hyaluronate-assocd. diseases)

L12 ANSWER 18 OF 20 HCAPLUS COPYRIGHT 2003 ACS on STN

ACCESSION NUMBER: 1981

1981:103813 HCAPLUS

DOCUMENT NUMBER:

94:103813

TITLE:

Photoreactive insulin derivatives: preparation and

characterization

AUTHOR(S):

Thamm, P.; Saunders, D.; Brandenburg, D.

CORPORATE SOURCE:

Deutsches Wollforschungsinst., Aachen, D-5100, Fed.

Rep. Ger.

SOURCE:

Insulin: Chem., Struct. Funct. Insulin Relat. Horm., Proc. Int. Insulin Symp., 2nd (1980), Meeting Date 1979, 309-16. Editor(s): Brandenburg, Dietrich; Wollmer, Axel. de Gruyter: Berlin, Fed. Rep. Ger.

CODEN: 44BTA8

DOCUMENT TYPE:

Conference English

LANGUAGE:

AB 1A-(R-Gly)-insulin (I) [R = 4,2-N3(O2N)C6H3], NB-(R-Gly)-29B-[N6-(R-Gly)-L-lysine]insulin, 2B,2B'bis(R1-L-Val)insulin-29B,29B'adipoyl dimer [R1 = 4,2-N3(O2N)C6H3CH2CO], NB-R1-insulin, 2B-(R1-L-Val)-1B-de-L-phenylalanineinsulin, and 29B-(N6-R1-L-Lys)insulin were prepd. Thus, NA-(Me3CO2C)-insulin was acylated by MeSO2CH2CH2O2COR2 (R2 = succinimido) and treated with F3CCO2H to give NB-(MeSO2CH2CH2O2C)-29B-[N6-(MeSO2CH2CH2O2C)-L-lysine]insulin, which underwent consecutive treatment with PhCSN, Edman degrdn., acylation by R-Gly-OR2, and deblocking to give I.

IT 76688-33-0P

L12 ANSWER 19 OF 20 HCAPLUS COPYRIGHT 2003 ACS on STN

ACCESSION NUMBER:

1979:421047 HCAPLUS

DOCUMENT NUMBER:

91:21047

TITLE:

Semisynthetic experiments in the C-terminal range of

insulin

AUTHOR(S):

Gattner, H. G.; Schmitt, E. W.; Naithani, V. K. Dtsch. Wollforschungsinst., Aachen, Fed. Rep. Ger.

CORPORATE SOURCE: SOURCE:

Semisynth. Pept. Proteins, Pap. Int. Meet. Protein Semisynth. (1978), Meeting Date 1977, 181-91. Editor(s): Offord, R. E.; Di Bello, C. Academic:

London, Engl. CODEN: 39MMAW

DOCUMENT TYPE:

Conference English

LANGUAGE:

Insulin hexamethyl ester (I) was cleaved with trypsin to give des-octapeptide-(B23-30) insulin pentamethyl ester, which was cleaved with carboxypeptidase B to give des-nonapeptide-(B22-30) insulin pentamethyl ester. Des-pentapeptide-(B26-30) insulin pentamethyl ester and des-alanine-(B30) insulin pentamethyl ester were also prepd. In the sapon. of I, asparaginimide formed at position A21; thus, difficulties may arise during sapon. steps in the semisynthesis of insulins. Porcine insulin was cleaved by pepsin to give des-pentapeptide-(B26-30) insulin, which was N-acylated with BOC-N3 (BOC = Me3CO2C) to give the NA,NB-di-BOC deriv. The latter was coupled with H-Tyr(CMe3)-Thr(CMe3)-Pro-Lys(BOC)-Thr(CMe3)-OCMe3 by dicyclohexylcarbodiimide/hydroxybenzotriazole to give the protected insulin, which was deblocked with CF3CO2H to give semisynthetic human insulin (porcine and human insulin differ only in residue B30).

#### 10 032330 Russel

IT 69913-74-2P RL: RCT (Reactant); SPN (Synthetic preparation); PREP (Preparation); RACT (Reactant or reagent) (prepn. and selective hydrolysis of) IT 69913-73-1P RL: RCT (Reactant); SPN (Synthetic preparation); PREP (Preparation); RACT (Reactant or reagent) (prepn. and selective hydrolysis of, with chymotrypsin) IT69913-72-0P RL: SPN (Synthetic preparation); PREP (Preparation) (prepn. of) L12 ANSWER 20 OF 20 HCAPLUS COPYRIGHT 2003 ACS on STN ACCESSION NUMBER: 1978:191425 HCAPLUS DOCUMENT NUMBER: 88:191425 TITLE: Preparation and application of N.alpha.-B1, N.epsilon.-B29-bis(tert-butyloxycarbonyl)insulin AUTHOR(S): Friesen, Heinz Juergen; Naithani, Vinod K.; Gattner, Hans Gregor CORPORATE SOURCE: Dtsch. Wollforschungsinst., Tech. Hochsch. Aachen, Aachen, Fed. Rep. Ger. SOURCE: Hoppe-Seyler's Zeitschrift fuer Physiologische Chemie (1978), 359(1), 103-11CODEN: HSZPAZ; ISSN: 0018-4888 DOCUMENT TYPE: Journal LANGUAGE: English The title compd. (I) was prepd. in 80-90% yield by acylating the .alpha.-N1 and .epsilon.-NB29 amino groups of NA1-trifluoroacetyl-insulin with BOCN3 (BOC = Me3CO2C) and cleaving the trifluoroacetyl group from the resulting triacylated insulin by NH4HCO3/NH3. I was also prepd. in 65% yield from NA1-citraconyl-insulin. The A1 glycine was cleaved from I by an Edman degrdn. and the resulting deriv. was BOC-deblocked with CF3CO2H to give des-GlyA1-insulin (II). NA1-Guanidinoacetyl-insulin (III) was prepd. from I. The in vitro biol. activities of I, II, and III are given. ΙT 66525-60-8P RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); SPN (Synthetic preparation); BIOL (Biological study); PREP (Preparation) (prepn. and biol. activity of) TT 66525-59-5P RL: RCT (Reactant); SPN (Synthetic preparation); PREP (Preparation); RACT (Reactant or reagent) (prepn. and reaction with trifluoroacetic acid) => select hit rn 112 1-20 E# OR SYSTEM LIMIT REACHED WHILE PROCESSING ANSWER 14 E1 THROUGH E999 ASSIGNED

=> fil req FILE 'REGISTRY' ENTERED AT 18:56:12 ON 28 JUL 2003 USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT. PLEASE SEE "HELP USAGETERMS" FOR DETAILS. COPYRIGHT (C) 2003 American Chemical Society (ACS)

Property values tagged with IC are from the ZIC/VINITI data file provided by InfoChem.

STRUCTURE FILE UPDATES: 27 JUL 2003 HIGHEST RN 556005-78-8 DICTIONARY FILE UPDATES: 27 JUL 2003 HIGHEST RN 556005-78-8

TSCA INFORMATION NOW CURRENT THROUGH JANUARY 6, 2003

Please note that search-term pricing does apply when conducting SmartSELECT searches.

Crossover limits have been increased. See HELP CROSSOVER for details.

Experimental and calculated property data are now available. See HELP PROPERTIES for more information. See STNote 27, Searching Properties in the CAS Registry File, for complete details: http://www.cas.org/ONLINE/STN/STNOTES/stnotes27.pdf

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(FILE 'HCAPLUS' ENTERED AT 18:42:39 ON 28 JUL 2003) SELECT HIT RN L12 1-20

FILE 'REGISTRY' ENTERED AT 18:43:30 ON 28 JUL 2003 L14 999 S E1-E999

FILE 'HCAPLUS' ENTERED AT 18:47:04 ON 28 JUL 2003 DEL SELECT SELECT HIT RN L12 14-20

FILE 'REGISTRY' ENTERED AT 18:47:32 ON 28 JUL 2003 L15 999 S E1-E999

FILE 'HCAPLUS' ENTERED AT 18:49:50 ON 28 JUL 2003 DEL SELECT Y SELECT HIT RN L12 15-20

FILE 'REGISTRY' ENTERED AT 18:50:12 ON 28 JUL 2003 L16 999 S E1-E999

FILE 'HCAPLUS' ENTERED AT 18:53:32 ON 28 JUL 2003 DEL SELECT Y SELECT HIT RN L12 17-20

FILE 'REGISTRY' ENTERED AT 18:53:59 ON 28 JUL 2003 L17 7 S E1-E7 L18 11 S (L15 OR L16 OR L17) AND L1

FILE 'HCAPLUS' ENTERED AT 18:54:49 ON 28 JUL 2003

FILE 'REGISTRY' ENTERED AT 18:56:12 ON 28 JUL 2003

=> d .seq 118 1-11

ANSWER 1 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN L18

402741-21-3 REGISTRY

CN L-Valine, glycyl-L-arginyl-L-arginyl-L-alpha.-aspartyl-L-phenylalanyl-Lprolyl-L-.alpha.-glutamyl-L-.alpha.-glutamyl-L-valyl-L-threonyl-Lisoleucyl-L-valyl-L-.alpha.-glutamyl-L-.alpha.-glutamyl-L-leucyl-L-arginyl-L-arginyl-L-arginyl-L-histidyl-L-alanyl-L-.alpha.-aspartyl- (9CI) (CA INDEX NAME)

OTHER NAMES:

Glucagon (Equus caballus gene GCG exon 4 fragment plus exon 5) CN SOL

RN 402741-21-3 REGISTRY

1 GRRDFPEEVT IVEELRRRHA DV SEQ

HITS AT: 11-14 \*\*RELATED SEQUENCES AVAILABLE WITH SEOLINK\*\* REFERENCE 1: 138:332694 REFERENCE 2: 136:211522 T.18 ANSWER 2 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN RN 400621-68-3 REGISTRY CN L-Serine, L-valyl-L-leucyl-L-alpha.-aspartyl-L-leucyl-L-valyl-Lphenylalanyl-L-seryl-L-.alpha.-glutamylglycylglycyl-L-isoleucyl-L-valyl-Lleucyl-L-seryl-L-phenylalanyl-L-arginyl-L-asparaginyl-L-leucyl-L-.alpha.glutamyl-L-arginyl-L-methionyl-L-valyl-L-leucyl-L-isoleucyl-L-leucyl-L-.alpha.-glutamyl-L-threonyl-L-histidyl-L-valyl- (9CI) (CA INDEX NAME) OTHER NAMES: CN1219: PN: WO0157272 SEQID: 28612 claimed sequence 1259: PN: WO0157278 SEQID: 20659 claimed CN CN 1763: PN: WO0157275 SEQID: 27737 claimed CN 2007: PN: WO0157274 SEQID: 22016 claimed sequence CN 2121: PN: WO0157273 SEQID: 28308 claimed 2276: PN: WO0157276 SEQID: 28320 claimed CN 3168: PN: WO0186003 SEQID: 27210 claimed CN 543: PN: US20020048763 SEQID: 35544 claimed CN Protein (human bone marrow clone WO0152276-SEQID-28320 exon-encoded CN fragment) Protein (human brain clone WO0157275-SEQID-27737 exon-encoded fragment) CN Protein (human BT474 cell clone WO0157271-SEQID-12620 exon-encoded CN CNProtein (human cervix cell clone WO0157278-SEQID-20659 exon-encoded fragment) Protein (human clone US20020048763-SEQID-35544 exon-encoded fragment) CN Protein (human clone WO01057273-SEQID-28308 exon-encoded fragment) CN Protein (human clone WO0157274-SEQID-22016 exon-encoded fragment) CN CN Protein (human fetal liver clone WO0157277-SEQID-27461 exon-encoded fragment) CN Protein (human lung clone WO0186003-SEQID-27210 exon-encoded fragment) CN Protein (human placenta clone WO0157272-SEQID-28612 exon-encoded fragment) SOL RN 400621-68-3 REGISTRY 1 VLDLVFSEGG IVLSFRNLER MVLILETHVS SEO HITS AT: 9 - 12\*\*RELATED SEQUENCES AVAILABLE WITH SEOLINK\*\* REFERENCE 1: 137:28982 REFERENCE 2: 137:16533 REFERENCE 136:351355 3: REFERENCE 4: 136:351347 REFERENCE 136:305084 5: REFERENCE 6: 136:211829 REFERENCE 7: 136:195269

8:

136:195264

REFERENCE

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REFERENCE
                136:195263
REFERENCE 10:
                136:178933
L18
    ANSWER 3 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
RN
     400620-15-7 REGISTRY
     L-Serine, L-.alpha.-aspartyl-L-phenylalanylglycylglycyl-L-histidyl-L-
CN
     histidyl-L-glutaminyl-L-leucylglycyl-L-prolylglycyl-L-leucyl-L-tryptophyl-
     L-threonyl-L-.alpha.-glutamylglycyl-L-cysteinyl-L-prolylglycyl-L-cysteinyl-
     L-valyl-L-histidyl-L-glutaminylglycyl-L-tyrosyl-L-glutaminyl-L-leucyl-L-
     prolyl-L-arginyl-L-leucyl-L-.alpha.-aspartyl- (9CI) (CA INDEX NAME)
OTHER NAMES:
CN
     1178: PN: WO0157275 SEQID: 27148 claimed
CN
     1447: PN: WO0157274 SEQID: 21456 claimed sequence
CN
     1554: PN: WO0157273 SEQID: 27737 claimed
CN
     1695: PN: WO0157276 SEQID: 27739 claimed
     2684: PN: WO0186003 SEQID: 26721 claimed
CN
     4941: PN: US20020048763 SEQID: 34984 claimed
CN
     600: PN: WO0157272 SEQID: 27993 claimed sequence
CN
CN
     685: PN: WO0157278 SEQID: 20085 claimed
CN
     Protein (human bone marrow clone WO0152276-SEQID-27739 exon-encoded
     Protein (human brain clone WO0157275-SEQID-27148 exon-encoded fragment)
CN
CN
     Protein (human BT474 cell clone WO0157271-SEQID-12057 exon-encoded
CN
     Protein (human cervix cell clone WO0157278-SEQID-20085 exon-encoded
     fragment)
CN
     Protein (human clone US20020048763-SEQID-34984 exon-encoded fragment)
CN
     Protein (human clone WO01057273-SEQID-27737 exon-encoded fragment)
CN
     Protein (human clone WO0157274-SEQID-21456 exon-encoded fragment)
CN
     Protein (human fetal liver clone WO0157277-SEQID-26885 exon-encoded
CN
     Protein (human lung clone WO0186003-SEQID-26721 exon-encoded fragment)
CN
     Protein (human placenta clone WO0157272-SEQID-27993 exon-encoded fragment)
SQL
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RN
     400620-15-7 REGISTRY
SEO
         1 DFGGHHQLGP GLWTEGCPGC VHQGYQLPRL DS
           25-28
HITS AT:
**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
REFERENCE
                137:28981
REFERENCE
            2:
                137:16533
REFERENCE
            3:
                136:351355
REFERENCE
                136:351347
            4:
REFERENCE
            5:
                136:305084
REFERENCE
            6:
                136:211829
            7:
                136:195269
REFERENCE
            8:
                136:195264
REFERENCE
REFERENCE
                136:195263
```

REFERENCE

10:

136:178933

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L18
      ANSWER 4 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
      400618-23-7 REGISTRY
 RN
      L-Asparagine, L-isoleucylglycylglycyl-L-isoleucyl-L-valylglycyl-L-
 CN
      methionyl-L-glutaminyl-L-leucyl-L-threonyl-L-.alpha.-glutamyl-L-leucyl-L-
      lysyl-L-threonyl-L-leucyl-L-leucyl-L-cysteinyl-L-valyl-L-alanyl-L-.alpha.-
      glutamylglycyl- (9CI) (CA INDEX NAME)
 OTHER NAMES:
      140: PN: WO0157272 SEQID: 27126 claimed sequence
 CN
      1862: PN: WO0186003 SEQID: 25888 claimed
 CN
 CN
      314: PN: WO0157275 SEQID: 26282 claimed
      4115: PN: US20020048763 SEQID: 34149 claimed
 CN
 CN
      612: PN: WO0157274 SEQID: 20621 claimed sequence
 CN
      716: PN: WO0157273 SEQID: 26887 claimed
CN
      833: PN: WO0157276 SEQID: 26877 claimed
CN
      878: PN: WO0157278 SEQID: 19270 claimed
CN
      Protein (human bone marrow clone WO0152276-SEQID-26877 exon-encoded
      fragment)
      Protein (human brain clone WO0157275-SEQID-26282 exon-encoded fragment)
CN
      Protein (human BT474 cell clone WO0157271-SEQID-11185 exon-encoded
CN
     Protein (human cervix cell clone WO0157278-SEQID-19270 exon-encoded
CN
      fragment)
     Protein (human clone US20020048763-SEQID-34149 exon-encoded fragment)
CN
     Protein (human clone WO01057273-SEQID-26887 exon-encoded fragment)
CN
     Protein (human clone WO0157274-SEQID-20621 exon-encoded fragment)
CN
CN
     Protein (human fetal liver clone WO0157277-SEQID-26027 exon-encoded
     Protein (human lung clone WO0186003-SEQID-25888 exon-encoded fragment)
CN
     Protein (human placenta clone WO0157272-SEQID-27126 exon-encoded fragment)
CN
SOL
RN
     400618-23-7 REGISTRY
SEQ
         1 IGGIVGMQLT ELKTLLCVAE GN
HITS AT:
           2-5
**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
REFERENCE
                137:28981
            1:
REFERENCE
            2:
                137:16533
REFERENCE
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                136:351355
REFERENCE
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                136:351347
REFERENCE
            5:
                136:305084
REFERENCE
            6:
                136:211829
REFERENCE
            7:
                136:195269
REFERENÇE
            8:
                136:195264
REFERENCE
            9:
                136:195263
REFERENCE 10:
                136:178933
    ANSWER 5 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
RN
    254965-50-9 REGISTRY
    L-Tyrosine, L-methionyl-L-alanyl-L-leucyl-L-glutaminyl-L-leucyl-L-prolyl-L-
CN
    tyrosyl- (9CI) (CA INDEX NAME)
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27: PN: WO0001841 PAGE: 42 claimed sequence

OTHER NAMES:

CN

SQL 254965-50-9 REGISTRY RN 1 MALQLPYY SEQ HITS AT: REFERENCE 1: 132:102856 L18 ANSWER 6 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN RN 76688-33-0 REGISTRY Insulin (cattle), NA-[N-(4-azido-2-nitrophenyl)glycyl]-NB-[N-(4-azido-2-CN nitrophenyl)glycyl]-29B-[N6-(trifluoroacetyl)-L-lysine]- (9CI) (CA INDEX NAME) OTHER CA INDEX NAMES: 3,4,44,45,90,91-Hexathia-8,11,14,17,20,23,26,29,32,35,38,41,48,51,54,57,60 ,63,66,69,72,75,78,81,84,86-hexacosaazabicyclo[72.11.7]dononacontane, cyclic peptide deriv. Insulin (ox), NA-[N-(4-azido-2-nitrophenyl)glycyl]-NB-[N-(4-azido-2-CN nitrophenyl)glycyl]-29B-[N6-(trifluoroacetyl)-L-lysine]-NTE multichain modified (modifications unspecified) ----- location ----- description Cys-8 - Cys-8' disulfide bridge Cys-20 - Cys-21' disulfide bridge Cys-7' - Cys-12' disulfide bridge bridge bridge bridge SQL 53,31,22 RN**76688-33-0** REGISTRY SEQ 1 GGIVEQCCAS VCSLYQLENY CN ==== 1 - 4HITS AT: REFERENCE 1: 94:103813 L18 ANSWER 7 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN 69913-74-2 REGISTRY RN (1A-21A), (1B-29B)-Insulin (human), hexamethyl ester (9CI) (CA INDEX NAME) CN OTHER CA INDEX NAMES: 3,4,44,45,90,91-Hexathia-8,11,14,17,20,23,26,29,32,35,38,41,48,51,54,57,60 ,63,66,69,72,75,78,81,84,86-hexacosaazabicyclo[72.11.7]dononacontane, cyclic peptide deriv. Insulin (ox), 8A-L-threonine-10A-L-isoleucine-30B-de-L-alanine-, CN hexamethyl ester NTE multichain modified (modifications unspecified) \_\_\_\_\_ ----- location ----- description Cys-7 - Cys-7' disulfide bridge Cys-19 - Cys-20' disulfide bridge Cys-6' - Cys-11' disulfide bridge bridge bridge bridge SQL 50,29,21 69913-74-2 REGISTRY RN

1 GIVEECCTSI CSLYQLENYC N

SEQ

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2-5
HITS AT:
REFERENCE
           1: 91:21047
L18 ANSWER 8 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
     69913-73-1 REGISTRY
RN
CN
     (1A-21A), (1B-25B)-Insulin (human), hexamethyl ester (9CI) (CA INDEX NAME)
OTHER CA INDEX NAMES:
CN
     3,4,44,45,90,91-Hexathia-8,11,14,17,20,23,26,29,32,35,38,41,48,51,54,57,60
     ,63,66,69,72,75,78,81,84,86-hexacosaazabicyclo[72.11.7]dononacontane,
     cyclic peptide deriv.
CN
     Insulin (ox), 8A-L-threonine-10A-L-isoleucine-26B-de-L-tyrosine-27B-de-L-
     threonine-28B-de-L-proline-29B-de-L-lysine-30B-de-L-alanine-, hexamethyl
    multichain
     modified (modifications unspecified).
_____
                ----- location ----- description
           Cys-7 - Cys-7' disulfide bridge
Cys-19 - Cys-20' disulfide bridge
Cys-6' - Cys-11' disulfide bridge
bridge
bridge
bridge
        SQL 46,25,21
    69913-73-1 REGISTRY
RN
SEQ
        1 GIVEECCTSI CSLYQLENYC N
HITS AT:
          2 - 5
**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
REFERENCE
           1: 91:21047
    ANSWER 9 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
L18
RN
     69913-72-0 REGISTRY
CN
     (1A-21A), (1B-25B)-Insulin (human), 4A,13B,17A,21A,21B-pentamethyl ester
     (9CI) (CA INDEX NAME)
OTHER CA INDEX NAMES:
    3, 4, 44, 45, 90, 91-Hexathia-8, 11, 14, 17, 20, 23, 26, 29, 32, 35, 38, 41, 48, 51, 54, 57, 60
     ,63,66,69,72,75,78,81,84,86-hexacosaazabicyclo[72.11.7]dononacontane,
    cyclic peptide deriv.
CN
    Insulin (ox), 8A-L-threonine-10A-L-isoleucine-26B-de-L-tyrosine-27B-de-L-
    threonine-28B-de-L-proline-29B-de-L-lysine-30B-de-L-alanine-,
    4A, 13B, 17A, 21A, 21B-pentamethyl ester
NTE
    multichain
    modified (modifications unspecified)
    ______
                ----- location ----- description
              Cys-7 - Cys-7' disulfide bridge
Cys-19 - Cys-20' disulfide bridge
Cys-6' - Cys-11' disulfide bridge
             Cys-7
bridge
bridge
bridge
SOL 46,25,21
    69913-72-0 REGISTRY
        1 GIVEECCTSI CSLYQLENYC N
SEQ
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HITS AT: 2 - 5

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

REFERENCE 1: 91:21047

L18 ANSWER 10 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN

RN 66525-60-8 REGISTRY

CN Insulin (cattle), NA-[N-(aminoiminomethyl)glycyl]- (9CI) (CA INDEX NAME) OTHER CA INDEX NAMES:

3,4,44,45,90,91-Hexathia-8,11,14,17,20,23,26,29,32,35,38,41,48,51,54,57,60 ,63,66,69,72,75,78,81,84,86-hexacosaazabicyclo[72.11.7]dononacontane, cyclic peptide deriv.

Insulin (ox), NA-[N-(aminoiminomethyl)glycyl]-CN

NTE multichain

modified (modifications unspecified)

type	lc	cation	description
bridge bridge bridge	Cys-7 Cys-19 Cys-7'	- Cys-8' - Cys-21' - Cys-12'	disulfide bridge disulfide bridge disulfide bridge
SQL 52,30,22			

RN 66525-60-8 REGISTRY

1 GGIVEQCCAS VCSLYQLENY CN SEO

HITS AT: 1 - 4

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

REFERENCE 1: 88:191425

L18 ANSWER 11 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN

RN 66525-59-5 REGISTRY

CN Insulin (cattle), NA-[N-(aminoiminomethyl)glycyl]-NB-[(1,1dimethylethoxy)carbonyl]-29B-[N6-[(1,1-dimethylethoxy)carbonyl]-L-lysine]-(9CI) (CA INDEX NAME)

OTHER CA INDEX NAMES:

3,4,44,45,90,91-Hexathia-8,11,14,17,20,23,26,29,32,35,38,41,48,51,54,57,60 CN ,63,66,69,72,75,78,81,84,86-hexacosaazabicyclo[72.11.7]dononacontane, cyclic peptide deriv.

CN Insulin (ox), NA-[N-(aminoiminomethyl)glycyl]-NB-[(1,1dimethylethoxy)carbonyl]-29B-[N6-[(1,1-dimethylethoxy)carbonyl]-L-lysine]-

NTE multichain

modified (modifications unspecified)

type	loca	tion	desci	ription	·
bridge bridge bridge	Cys-7 Cys-19 Cys-7'	- Cys-8' - Cys-21' - Cys-12'	disulfide disulfide disulfide	bridge	·

SOL 52,30,22

66525-59-5 REGISTRY

1 GGIVEQCCAS VCSLYQLENY CN SEQ

HITS AT:

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

REFERENCE 1: 88:191425

=>

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FILE COVERS 1907 - 28 Jul 2003 VOL 139 ISS 5 FILE LAST UPDATED: 27 Jul 2003 (20030727/ED)

This file contains CAS Registry Numbers for easy and accurate substance identification.

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=>
=> d stat que 113
          13173 SEA FILE=REGISTRY ABB=ON PLU=ON GIVE/SQSP
L2
L4
          50431 SEA FILE=REGISTRY ABB=ON
                                         PLU=ON
                                                 BONE OR FATTY (W) ACID? OR
                ACYLAT?
          10042 SEA FILE=REGISTRY ABB=ON PLU=ON L2 NOT INSULIN?
L5
         686265 SEA FILE=HCAPLUS ABB=ON PLU=ON L4 OR BONE OR FATTY(W)ACID?
Ь7
                OR ACYLAT? OR OSTEOPOROS?
            252 SEA FILE=REGISTRY ABB=ON PLU=ON L5 AND SQL<=25
L9
L11
            142 SEA FILE=HCAPLUS ABB=ON PLU=ON L9
L13
              4 SEA FILE=HCAPLUS ABB=ON PLU=ON L11 AND L7
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=> d ibib abs hitrn 113 1-4

=> =>

L13 ANSWER 1 OF 4 HCAPLUS COPYRIGHT 2003 ACS on STN

ACCESSION NUMBER: 2002:850307 HCAPLUS

DOCUMENT NUMBER: 137:346244

TITLE: Tissue remodeling with compds. comprising a sequence

from TGF-.beta. super family Ser/Thr/kinase receptors

INVENTOR(S): Ben-Sasson, Shmuel

PATENT ASSIGNEE(S): Children's Medical Center Corporation, USA

SOURCE: U.S. Pat. Appl. Publ., 51 pp., Cont.-in-part of Appl.

No. PCT/US00/32852.

CODEN: USXXCO

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 3

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
				- <b></b>
US 2002165150	A1	20021107	US 2001-32330	20011231

```
A2
                               20010614
                                               WO 2000-US32852 20001204
     WO 2001042280
     WO 2001042280
                         A3
                               20020307
              AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN,
              CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT,
              LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU,
              SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN,
              YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
          RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,
              DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF,
              BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
PRIORITY APPLN. INFO.:
                                            US 1998-161094
                                                              B2 19980925
                                            WO 2000-US32851
                                                              W 20001204
                                            WO 2000-US32852 A2 20001204
                                            US 1999-458491
                                                              A1 19991209
AB
     The invention concerns a method for the modulation of tissue-remodeling
     processes, by contacting the tissue to be remodeled with a compd.
     comprising a sequence derived from certain regions of TGF-.beta. super
     family Ser/Thr/kinase receptors.
     332350-87-5, BMP receptor kinase-2
IT
     RL: BSU (Biological study, unclassified); BIOL (Biological study)
         (tissue remodeling with compds. comprising a sequence from TGF-.beta.
        super family Ser/Thr/kinase receptors)
     474526-71-1 474526-72-2 474526-73-3
ΙT
     474526-74-4 474526-78-8 474526-96-0
     RL: PAC (Pharmacological activity); THU (Therapeutic use); BIOL
     (Biological study); USES (Uses)
         (tissue remodeling with compds. comprising a sequence from TGF-.beta.
        super family Ser/Thr/kinase receptors)
L13 ANSWER 2 OF 4 HCAPLUS COPYRIGHT 2003 ACS on STN
                           2000:513533 HCAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                           133:140232
TITLE:
                           Monodisperse hexameric acylated insulin
                           analog formulations
                           Ng, Kingman; Li, Shun; Watts, Eric Alan
INVENTOR(S):
PATENT ASSIGNEE(S):
                           Eli Lilly and Company, USA
SOURCE:
                           PCT Int. Appl., 56 pp.
                           CODEN: PIXXD2
DOCUMENT TYPE:
                           Patent
LANGUAGE:
                           English
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
     PATENT NO.
                        KIND
                              DATE
                                               APPLICATION NO.
                                                                  DATE
                        ____
                                               _____
     WO 2000043034
                        A2
                               20000727
                                               WO 2000-US1627
                                                                  20000126
     WO 2000043034
                        A3
                               20001228
             AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD,
              MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK,
```

Т3 20030216 ES 2000-904496 20000126 ES 2180511 PRIORITY APPLN. INFO.: US 1999-117291P P 19990126 W WO 2000-US1627 20000126 The present invention provides formulations and methods for prepg. AΒ formulations contg. an aq. soln. at a pH of greater than about 7.9. The aq. soln. includes an isotonicity agent, a phenolic deriv., zinc ions, and an acylated human insulin analog. More particularly, the invention relates to formulations having a pH of greater than about 7.9 that include a monoacylated human insulin analog such as an acylated des(B30) human insulin analog or an analog that comprises a native or modified human insulin A chain optionally modified at position A21, and a modified native human insulin B chain optionally modified at position B3 and modified at position B28, or at both positions B28 and B29, and that contains a lysine residue at either position B28 or B29 acylated with a fatty acid residue. The invention also provides a method for treating a patient suffering from hyperglycemia using the pharmaceutical formulations of the invention. 286410-19-3P ITRL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); PNU (Preparation, unclassified); RCT (Reactant); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); RACT (Reactant or reagent); USES (Uses) (monodisperse hexameric acylated insulin analog formulations) L13 ANSWER 3 OF 4 HCAPLUS COPYRIGHT 2003 ACS on STN 2000:10612 HCAPLUS ACCESSION NUMBER: DOCUMENT NUMBER: 132:73648 Lipophilic insulin derivatives soluble at TITLE: physiological pH with prolonged serum half-lives and biological activity Havelund, Svend; Halstrom, John; Jonassen, Ib; INVENTOR(S): Andersen, Asser Sloth; Markussen, Jan Novo Nordisk A/S, Den. PATENT ASSIGNEE(S): SOURCE: U.S., 47 pp., Cont.-in-part of U.S. 5,750,497. CODEN: USXXAM DOCUMENT TYPE: Patent LANGUAGE: English FAMILY ACC. NUM. COUNT: PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO. DATE
US 6011007	A	20000104	US 1997-975365 19971120
ZA 9407187	A	19950317	ZA 1994-7187 19940916
JP 2000060556	A2	20000229	JP 1999-221632 19940916
EP 1132404	A2	20010912	EP 2001-112992 19940916
EP 1132404	A3	20020327	
R: AT, BE,	CH, DE	, DK, ES,	FR, GB, GR, IT, LI, LU, NL, SE, PT, IE,
SI, LT			
JP 2002308899	A2	20021023	JP 2001-385921 19940916
US 5750497	A	19980512	US 1995-400256 19950308
AU 745983	B2	20020411	AU 2000-51960 20000811
PRIORITY APPLN. INFO	.:		DK 1993-1044 A 19930917
			US 1995-400256 A2 19950308
	•		US 1994-190829 A 19940202
			EP 1994-926816 A3 19940916
		•	JP 1995-508923 A3 19940916
			JP 1999-221632 A3 19940916

OTHER SOURCE(S): MARPAT 132:73648

AB Human insulin derivs. with improved soly. at physiol. pH and that retain biol. activity for longer than wild-type human insulin are described. The insulins are substituted at positions A21 and B3 with either being any amino acid except lysine, arginine, or cysteine. The phenylalanine at B1

may be deleted and the amino acid at position B30 may be deleted or substituted by any amino acid except lysine, arginine, or cysteine or by another amino acid that is lipophilic having a C10-24 side chain. If B30 is deleted or substituted, lysineB29 is modified by a carboxylic acid connected to the .epsilon.-amino group. When B30 is threonine or alanine and A21 and B3 are both asparagine, and phenylalanineB1 is present, then the insulin deriv. is always present as a Zn2 complex.

IT . 253597-47-6

RL: PRP (Properties)

(unclaimed protein sequence; lipophilic insulin derivs. sol. at physiol. pH with prolonged serum half-lives and biol. activity)

REFERENCE COUNT:

THERE ARE 25 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L13 ANSWER 4 OF 4 HCAPLUS COPYRIGHT 2003 ACS on STN

ACCESSION NUMBER:

1991:102773 HCAPLUS

DOCUMENT NUMBER:

114:102773

TITLE:

Studies on the total synthesis of an

A7, B7-dicarbainsulin III. Assembly of segments and

generation of biological activity

AUTHOR(S):

Videnov, G.; Buettner, Klaus; Casaretto, M.; Fohles,

Josef; Gattner, Hans Gregor; Stoev, S.; Brandenburg,

Dietrich

CORPORATE SOURCE:

Inst. Mol. Biol., Sofia, 1113, Bulg.

SOURCE:

Biological Chemistry Hoppe-Seyler (1990), 371(11),

1057-66

CODEN: BCHSEI; ISSN: 0177-3593

DOCUMENT TYPE: LANGUAGE:

Journal English

As a further contribution to the synthesis of an insulin analog with a stable A7-B7 interchain bond, the synthesis of A(8-21) by soln. methods, and of B(9-25) as well as [7-(2,7-diaminosuberic acid)]B(1-8) by solid phase methods is described. In the latter compd., the amino group of the diaminosuberic acid residue was acylated with A(1-6), and the resulting "U-peptide" sequentially elongated with the C-terminal A- and finally B-chain sequences. The conversion of the product into the disulfide moiety gave a mixt. which could not be resolved by currently available methods. However, the low biol. activity of the crude product indicates that the A7-B7 disulfide bond is not crucially important for the activity of insulin.

IT 126758-97-2

RL: RCT (Reactant); RACT (Reactant or reagent)

(peptide coupling of, with polymer-bound dicarbainsulin fragment)

IT 132167-71-6P

RL: RCT (Reactant); SPN (Synthetic preparation); PREP (Preparation); RACT (Reactant or reagent)

(prepn. and peptide coupling of, with insulin A-chain fragment)

IT 132167-76-1P

RL: SPN (Synthetic preparation); PREP (Preparation) (prepn. and protection of hydrazide group of)

=> select hit rn 113 1-4 E8 THROUGH E19 ASSIGNED

=> fil reg

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STRUCTURE FILE UPDATES:
                           27 JUL 2003
                                       HIGHEST RN 556005-78-8
DICTIONARY FILE UPDATES:
                          27 JUL 2003
                                      HIGHEST RN 556005-78-8
TSCA INFORMATION NOW CURRENT THROUGH JANUARY 6, 2003
  Please note that search-term pricing does apply when
  conducting SmartSELECT searches.
Crossover limits have been increased. See HELP CROSSOVER for details.
Experimental and calculated property data are now available. See HELP
PROPERTIES for more information. See STNote 27, Searching Properties
in the CAS Registry File, for complete details:
http://www.cas.org/ONLINE/STN/STNOTES/stnotes27.pdf
=>
=> s e8-e19
             1 126758-97-2/BI
                 (126758-97-2/RN)
             1 132167-71-6/BI
                 (132167-71-6/RN)
             1 132167-76-1/BI
                 (132167-76-1/RN)
             1 253597-47-6/BI
                 (253597-47-6/RN)
             1 286410-19-3/BI
                 (286410-19-3/RN)
             1 332350-87-5/BI
                 (332350-87-5/RN)
             1 474526-71-1/BI
                 (474526-71-1/RN)
             1 474526-72-2/BI
                 (474526-72-2/RN)
             1 474526-73-3/BI
                 (474526-73-3/RN)
             1 474526-74-4/BI
                 (474526-74-4/RN)
             1 474526-78-8/BI
                 (474526-78-8/RN)
             1 474526-96-0/BI
                 (474526-96-0/RN)
L19
            12 (126758-97-2/BI OR 132167-71-6/BI OR 132167-76-1/BI OR 253597-47
               -6/BI OR 286410-19-3/BI OR 332350-87-5/BI OR 474526-71-1/BI OR
               474526-72-2/BI OR 474526-73-3/BI OR 474526-74-4/BI OR 474526-78-
               8/BI OR 474526-96-0/BI)
=> s 119 and 12
            11 L19 AND L2
L20
=> d .seq 120 1-11
L20
    ANSWER 1 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
     474526-96-0 REGISTRY
RN
     L-Tyrosine, N-(1-oxotetradecyl)qlycylqlycyl-L-isoleucyl-L-valyl-L-alpha.-
     glutamyl-L-.alpha.-glutamyl-L-tyrosyl-L-glutaminyl-L-leucyl-L-prolyl-
     (9CI) (CA INDEX NAME)
OTHER NAMES:
     21: PN: US20020165150 SEQID: 21 claimed protein
```

NTE modified (modifications unspecified)

```
----- location -----
                                                                                    description
  type
 ------
modification Gly-1 - undetermined modification
SQL 11
RN
         474526-96-0 REGISTRY
SEO
                  1 GGIVEEYQLP Y
                      ==== ...
HITS AT:
                     2-5
 **RELATED SEQUENCES AVAILABLE WITH SEQLINK**
REFERENCE
                    1: 137:346244
L20 ANSWER 2 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
RN
          474526-78-8 REGISTRY
          \verb|L-Phenylalanine|, N-(1-oxooctadecyl)glycylglycyl-L-isoleucyl-L-valyl-L-isoleucyl-L-valyl-L-isoleucyl-L-valyl-L-isoleucyl-L-valyl-L-isoleucyl-L-valyl-L-isoleucyl-L-valyl-L-isoleucyl-L-valyl-L-isoleucyl-L-valyl-L-isoleucyl-L-isoleucyl-L-valyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L-isoleucyl-L
          .alpha.-glutamyl-L-.alpha.-aspartyl-L-tyrosyl-L-arginyl-L-prolyl-L-prolyl-
          (9CI) (CA INDEX NAME)
OTHER NAMES:
       10: PN: US20020165150 SEQID: 10 claimed protein
NTE modified (modifications unspecified)
______
 type ----- location ----- description
 modification Gly-1 - -
                                                                         undetermined modification
SOL 11
         474526-78-8 REGISTRY
RN
SEO
                  1 GGIVEDYRPP F
HITS AT:
                     2-5
**RELATED SEOUENCES AVAILABLE WITH SEOLINK**
REFERENCE
                    1: 137:346244
L20 ANSWER 3 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
RN
          474526-74-4 REGISTRY
CN
         L-Proline, N-(1-oxotetradecyl)glycyl-L-valyl-L-asparaginylglycyl-L-
         isoleucyl-L-valyl-L-.alpha.-glutamyl-L-.alpha.-aspartyl-L-tyrosyl-L-arginyl- (9CI) (CA INDEX NAME)
OTHER NAMES:
CN 6: PN: US20020165150 SEQID: 6 claimed protein
NTE modified (modifications unspecified)
_____
                                ----- location ----- description
modification Gly-1 - undetermined modification
SOL 11
RN
         474526-74-4 REGISTRY
                 1 GVNGIVEDYR P
SEO
HITS AT:
                     4 - 7
REFERENCE 1: 137:346244
L20 ANSWER 4 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
          474526-73-3 REGISTRY
RN
```

```
CN
     L-Proline, N-(1-oxotetradecyl)glycylglycyl-L-isoleucyl-L-valyl-L-.alpha.-
     glutamyl-L-.alpha.-aspartyl-L-tyrosyl-L-arginyl- (9CI) (CA INDEX NAME)
OTHER NAMES:
    5: PN: US20020165150 SEQID: 5 claimed protein
CN
NTE modified (modifications unspecified)
          ----- location ----- description
modification Gly-1 -
                                         undetermined modification
SQL 9
    474526-73-3 REGISTRY
RN
        1 GGIVEDYRP
SEQ
HITS AT:
          2-5
REFERENCE 1: 137:346244
L20 ANSWER 5 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
RN
    474526-72-2 REGISTRY
CN
    L-Prolinamide, N-(1-oxotetradecyl)glycylglycyl-L-isoleucyl-L-valyl-L-
     .alpha.-glutamyl-L-.alpha.-aspartyl-L-tyrosyl-L-arginyl-L-prolyl- (9CI)
     (CA INDEX NAME)
OTHER NAMES:
    4: PN: US20020165150 SEQID: 4 claimed protein
NTE modified
                ----- location -----
terminal mod. Pro-10 modification Gly-1
                                  C-terminal amide undetermined modification
SOL 10
    474526-72-2 REGISTRY
RN
SEQ
        1 · GGIVEDYRPP
HITS AT:
          2-5
          1: 137:346244
REFERENCE
    ANSWER 6 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
L20
    474526-71-1 REGISTRY
RN
    L-Phenylalanine, N-(1-oxotetradecyl)glycylglycyl-L-isoleucyl-L-valyl-L-
CN
     .alpha.-glutamyl-L-.alpha.-aspartyl-L-tyrosyl-L-arginyl-L-prolyl-L-prolyl-
     (9CI) (CA INDEX NAME)
OTHER NAMES:
CN 3: PN: US20020165150 SEQID: 3 claimed protein
NTE modified (modifications unspecified)
                ----- location ----- description
type
modification Gly-1
                                    undetermined modification
SOL 11
RN
    474526-71-1 REGISTRY
SEO
        1 GGIVEDYRPP F
HITS AT:
          2-5
**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
```

```
REFERENCE 1: 137:346244
   ANSWER 7 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
T<sub>2</sub>0
RN
   286410-19-3 REGISTRY
   Peptide, (Gly-Ile-Val-Glu-Gln-Cys-Cys-Thr-Ser-Ile-Cys-Ser-Leu-Tyr-Gln-Leu-
CN
   Glu-Asn-Tyr-Cys-Xaa-Xaa) (9CI) (CA INDEX NAME)
OTHER NAMES:
CN 1: PN: WO0043512 SEQID: 1 claimed protein
NTE
             ----- location ----- description
uncommon Aaa-21
uncommon Aaa-22
          _____
SQL 22
   286410-19-3 REGISTRY
RN
       1 GIVEQCCTSI CSLYQLENYC XX
SEQ
HITS AT:
        1 - 4
REFERENCE 1: 133:140232
L20 ANSWER 8 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
   253597-47-6 REGISTRY
RN
   1: PN: US6011007 SEQID: 1 unclaimed protein (9CI) (CA INDEX NAME)
CN
NTE
      _____
            ----- location -----
_____
uncommon Aaa-21 -
SQL 21
   253597-47-6 REGISTRY
RN
       1 GIVEQCCTSI CSLYQLENYC X
SEO
HITS AT: 1-4
REFERENCE 1: 132:73648
L20 ANSWER 9 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN
    132167-76-1 REGISTRY
RN
    Glycine, N-[(phenylmethoxy)carbonyl]-L-phenylalanyl-L-valyl-L-asparaginyl-
CN
    L-glutaminyl-L-histidyl-L-leucyl-7-carboxy-N7-[N-[N2-[N-[N-[N-[N-[(1,1-
    dimethylethoxy)carbonyl]glycyl]-L-isoleucyl]-L-valyl]-L-.alpha.-glutamyl]-
    L-glutaminyl]-3-[(1,1-dimethylethyl)dithio]-L-alanyl]-L-2,7-
    diaminoheptanoyl-, 7(5)-(1,1-dimethylethyl) ester, 8-hydrazide (9CI) (CA
    INDEX NAME)
NTE multichain
   modified (modifications unspecified)
            ----- location ----- description
bridge Dsu-7 - Leu-6' amide bridge uncommon Dsu-7 - -
_____
SQL 14,8,6
RN 132167-76-1 REGISTRY
      1 GIVEOCXG
SEQ
```

HITS AT: 1 - 4\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\* 1: 114:102773 REFERENCE ANSWER 10 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN L20 RN 132167-71-6 REGISTRY Glycine, N-[(phenylmethoxy)carbonyl]-L-phenylalanyl-L-valyl-L-asparaginyl-CN dimethylethoxy)carbonyl]glycyl]-L-isoleucyl]-L-valyl]-L-.alpha.-glutamyl]-L-glutaminyl]-3-[(1,1-dimethylethyl)dithio]-L-alanyl]-L-2,7diaminoheptanoyl-, 7(5)-(1,1-dimethylethyl) ester, 8-[2-[[2-[[4-(methylsulfonyl)phenyl]sulfonyl]ethoxy]carbonyl]hydrazide] (9CI) INDEX NAME) NTE multichain modified (modifications unspecified) \_\_\_\_\_ ----- location ----- description bridge Dsu-7 - Leu-6' amide bridge uncommon Dsu-7 - -SQL 14,8,6 RN **132167-71-6** REGISTRY 1 GIVEOCXG SEQ HITS AT: 1 - 4\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\* 1: 114:102773 REFERENCE L20 ANSWER 11 OF 11 REGISTRY COPYRIGHT 2003 ACS on STN **126758-97-2** REGISTRY RNCN isoleucyl]-L-valyl]-L-.alpha.-glutamyl]-L-glutaminyl]-3-[(1,1dimethylethyl)dithio]-, 5-(1,1-dimethylethyl) ester (9CI) (CA INDEX NAME) NTE modified (modifications unspecified) type ----- location ----- description modification Gly-1 - (1,1-dimethylethoxy) can modification Glu-4 - 1,1-dimethylethyl< t-Bu> modification Cys-6 - (1,1-dimethylethyl)thio(1,1-dimethylethoxy) carbonyl<Boc> \_\_\_\_\_\_ SQL 6 **126758-97-2** REGISTRY RN 1 GIVEQC SEO HITS AT:

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

REFERENCE 1: 114:102773

REFERENCE 2: 112:199075